



SEQUENCE LISTING

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<120> NOVEL ANDROGEN-INDUCED SUPPRESSOR OF CELL PROLIFERATION AND USES THEREOF

<130> 34724-026

<140> US 09/512,581

<141> 2000-02-24

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<151> 1999-02-24

<160> 72

<170> PatentIn version 3.3

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 Phe Tyr Leu Leu Glu Asn Ile Ala Trp Val Lys Ser Tyr Asn Ile Cys
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Leu Phe Ser Val Ile Asn Asn Gly His Asn Gln Lys Val His Met His
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Gln Glu Leu Leu Asp Thr Val Leu Val Asn Leu Val Pro Ala His Lys
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Thr Ala Gln Ala Ile Glu Pro Tyr Ile Thr Thr Phe Phe Asn Gln Val
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Arg Leu Gln Val Val Lys Leu Leu Ala Lys Met Phe Gly Ala Lys Asp
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Arg Phe Asn Asp Ile His Val Pro Ile Arg Leu Glu Cys Val Lys Phe
325 330 335
Ala Ser His Cys Leu Met Asn His Pro Asp Leu Ala Lys Asp Leu Thr
340 345 350
Glu Tyr Leu Lys Val Arg Ser His Asp Pro Glu Glu Ala Ile Arg His
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Asp Val Ile Val Ser Ile Val Thr Ala Ala Lys Lys Asp Ile Leu Leu
370 375 380
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385 390 395 400
Arg Trp Arg Val Arg Lys Glu Ala Met Met Gly Leu Ala Gln Ile Tyr
405 410 415
Lys Lys Tyr Ala Leu Gln Ser Ala Ala Gly Lys Asp Ala Ala Lys Gln
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Ile Ala Trp Ile Lys Asp Lys Leu Leu His Ile Tyr Tyr Gln Asn Ser
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Glu	Gly	Cys	Val	Arg	Glu	Ile	Thr	Lys	Lys	Leu	Gly	Asn	Pro	Lys	Gln	580	585	590
Pro	Thr	Asn	Pro	Phe	Leu	Glu	Met	Ile	Lys	Phe	Leu	Leu	Glu	Arg	Ile	595	600	605
Ala	Pro	Val	His	Ile	Asp	Thr	Glu	Ser	Ile	Ser	Ala	Leu	Ile	Lys	Gln	610	615	620
Val	Asn	Lys	Ser	Ile	Asp	Gly	Thr	Ala	Asp	Asp	Glu	Asp	Glu	Gly	Val	625	630	635
Pro	Thr	Asp	Gln	Ala	Ile	Arg	Ala	Gly	Leu	Glu	Leu	Leu	Lys	Val	Leu	645	650	655
Ser	Phe	Thr	His	Pro	Ile	Ser	Phe	His	Ser	Ala	Glu	Thr	Phe	Glu	Ser	660	665	670
Leu	Leu	Ala	Cys	Leu	Lys	Met	Asp	Asp	Glu	Lys	Val	Ala	Glu	Ala	Ala	675	680	685
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His	Ile	Arg	Ser	Ala	Leu	Leu	Pro	Val	Leu	His	His	Lys	Ser	Lys	Lys	705	710	715
Gly	Pro	Pro	Arg	Gln	Ala	Lys	Tyr	Ala	Ile	His	Cys	Ile	His	Ala	Ile	725	730	735
Phe	Ser	Ser	Lys	Glu	Thr	Gln	Phe	Ala	Gln	Ile	Phe	Glu	Pro	Leu	His	740	745	750

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Arg Arg Leu Lys Met Val Val Lys Thr Phe Met Asp Met Asp Gln Asp
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Ser Glu Glu Glu Lys Glu Leu Tyr Leu Asn Leu Ala Leu His Leu Ala
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Ser Asp Phe Phe Leu Lys His Pro Gly Lys Asp Val Arg Leu Leu Val
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Ala Cys Cys Leu Ala Asp Ile Phe Arg Ile Tyr Ala Pro Glu Ala Pro
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Tyr Thr Ser Pro Asp Lys Leu Lys Asp Ile Phe Met Phe Ile Thr Arg
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Gln Leu Lys Gly Leu Glu Asp Thr Lys Ser Pro Gln Phe Asn Arg Tyr
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Ala Ser His Cys Leu Met Asn His Pro Asp Leu Ala Lys Asp Leu Thr	
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Pro Thr Asp Gln Ala Ile Arg Ala Gly Leu Glu Leu Leu Lys Val Leu	
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Ile Thr Thr Phe Phe Asn Gln Val Leu Met Leu Gly Lys Thr Ser Ile	
235 240 245	
agc gat ttg tca gag cat gtc ttt gac tta att ttg gag ctc tac aat	941
Ser Asp Leu Ser Glu His Val Phe Asp Leu Ile Leu Glu Leu Tyr Asn	
250 255 260	

att gat agt cat ttg ctg ctc tct gtt tta ccc cag ctt gaa ttt aaa	989
Ile Asp Ser His Leu Leu Leu Ser Val Leu Pro Gln Leu Glu Phe Lys	
265 270 275 280	
tta aag agc aat gat aat gag gag cgc cta caa gtt gtt aaa cta ctg	1037
Leu Lys Ser Asn Asp Asn Glu Glu Arg Leu Gln Val Val Lys Leu Leu	
285 290 295	
gca aaa atg ttt ggg gca aag gat tca gaa ttg gct tct caa aac aag	1085
Ala Lys Met Phe Gly Ala Lys Asp Ser Glu Leu Ala Ser Gln Asn Lys	
300 305 310	
cca ctt tgg cag tgc tac ttg ggc agg ttt aat gat atc cat gta cca	1133
Pro Leu Trp Gln Cys Tyr Leu Gly Arg Phe Asn Asp Ile His Val Pro	
315 320 325	
atc cgc ctg gaa tgt gtg aaa ttt gct agc cat tgt ctc atg aac cat	1181
Ile Arg Leu Glu Cys Val Lys Phe Ala Ser His Cys Leu Met Asn His	
330 335 340	
cct gat tta gca aaa gac tta aca gag tat ctt aaa gtg agg tca cat	1229
Pro Asp Leu Ala Lys Asp Leu Thr Glu Tyr Leu Lys Val Arg Ser His	
345 350 355 360	
gac cct gag gaa gct att aga cat gat gtt att gtg tca ata gtt aca	1277
Asp Pro Glu Glu Ala Ile Arg His Asp Val Ile Val Ser Ile Val Thr	
365 370 375	
gct gct aaa aag gat att ctt ctg gtc aat gat cac tta ctt aat ttt	1325
Ala Ala Lys Lys Asp Ile Leu Leu Val Asn Asp His Leu Leu Asn Phe	
380 385 390	
gtg aga gag aga aca tta gac aaa cga tgg aga gta cgc aaa gaa gcc	1373
Val Arg Glu Arg Thr Leu Asp Lys Arg Trp Arg Val Arg Lys Glu Ala	
395 400 405	
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Met Met Gly Leu Ala Gln Ile Tyr Lys Lys Tyr Ala Leu Gln Ser Ala	
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gct gga aaa gat gct gca aaa cag ata gca tgg atc aaa gac aaa ttg	1469
Ala Gly Lys Asp Ala Ala Lys Gln Ile Ala Trp Ile Lys Asp Lys Leu	
425 430 435 440	
cta cat ata tat tat caa aat agt att gat gat cga cta ctt gtt gaa	1517
Leu His Ile Tyr Tyr Gln Asn Ser Ile Asp Asp Arg Leu Leu Val Glu	
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cgg atc ttt gct caa tac atg gtt cct cac aat tta gaa act aca gaa	1565
Arg Ile Phe Ala Gln Tyr Met Val Pro His Asn Leu Glu Thr Thr Glu	
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cgg atg aaa tgc tta tat tac ttg tat gcc aca ctg gat tta aat gct	1613
Arg Met Lys Cys Leu Tyr Tyr Leu Tyr Ala Thr Leu Asp Leu Asn Ala	
475 480 485	

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Val Lys Ala Leu Asn Glu Met Trp Lys Cys Gln Asn Leu Leu Arg His	
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Gln Val Lys Asp Leu Leu Asp Leu Ile Lys Gln Pro Lys Thr Asp Ala	
505 510 515 520	
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Ser Val Lys Ala Ile Phe Ser Lys Val Met Val Ile Thr Arg Asn Leu	
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Pro Asp Pro Gly Lys Ala Gln Asp Phe Met Lys Lys Phe Thr Gln Val	
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Leu Glu Asp Asp Glu Lys Ile Arg Lys Gln Leu Glu Val Leu Val Ser	
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Pro Thr Cys Ser Cys Lys Gln Ala Glu Gly Cys Val Arg Glu Ile Thr	
570 575 580	
aag aag ttg ggc aac ccc aaa cag cct aca aat cct ttc ctg gaa atg	1949
Lys Lys Leu Gly Asn Pro Lys Gln Pro Thr Asn Pro Phe Leu Glu Met	
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Ile Lys Phe Leu Leu Glu Arg Ile Ala Pro Val His Ile Asp Thr Glu	
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tct atc agt gct ctt att aaa caa gtg aac aaa tca ata gat gga aca	2045
Ser Ile Ser Ala Leu Ile Lys Gln Val Asn Lys Ser Ile Asp Gly Thr	
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Ala Asp Asp Glu Asp Glu Gly Val Pro Thr Asp Gln Ala Ile Arg Ala	
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Gly Leu Glu Leu Leu Lys Val Leu Ser Phe Thr His Pro Ile Ser Phe	
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Ser Lys Ile Glu Glu Asp Phe Pro His Ile Arg Ser Ala Leu Leu Pro	
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Val Leu His His Lys Ser Lys Lys Gly Pro Pro Arg Gln Ala Lys Tyr	
715 720 725	
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Ala Ile His Cys Ile His Ala Ile Phe Ser Ser Lys Glu Thr Gln Phe	
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Ala Gln Ile Phe Glu Pro Leu His Lys Ser Leu Asp Pro Ser Asn Leu	
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gaa cat ctc ata aca cca ttg gtt act att ggt cat att gct ctc ctt	2477
Glu His Leu Ile Thr Pro Leu Val Thr Ile Gly His Ile Ala Leu Leu	
765 770 775	
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Thr Lys Leu Trp Val Pro Asp Glu Glu Val Ser Pro Glu Thr Met Val	
810 815 820	
aaa att cag gct att aaa atg atg gtt cga tgg cta ctt gga atg aaa	2669
Lys Ile Gln Ala Ile Lys Met Met Val Arg Trp Leu Leu Gly Met Lys	
825 830 835 840	
aat aat cac agt aaa tca gga act tct acc tta aga ttg cta aca aca	2717
Asn Asn His Ser Lys Ser Gly Thr Ser Thr Leu Arg Leu Leu Thr Thr	
845 850 855	
ata ttg cat agt gat gga gac ttg aca gaa cag ggg aaa att agt aaa	2765
Ile Leu His Ser Asp Gly Asp Leu Thr Glu Gln Gly Lys Ile Ser Lys	
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Pro Asp Met Ser Arg Leu Arg Leu Ala Ala Gly Ser Ala Ile Val Lys	
875 880 885	
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Leu Ala Gln Glu Pro Cys Tyr His Glu Ile Ile Thr Leu Glu Gln Tyr	
890 895 900	
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Gln Leu Cys Ala Leu Ala Ile Asn Asp Glu Cys Tyr Gln Val Arg Gln	
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gtg ttt gcc cag aaa ctt cac aaa ggc ctt tcc cgt tta cgg ctt cca	2957
Val Phe Ala Gln Lys Leu His Lys Gly Leu Ser Arg Leu Arg Leu Pro	
925 930 935	

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Leu Glu Tyr Met Ala Ile Cys Ala Leu Cys Ala Lys Asp Pro Val Lys	
940 945 950	
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Glu Arg Arg Ala His Ala Arg Gln Cys Leu Val Lys Asn Ile Asn Val	
955 960 965	
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Arg Arg Glu Tyr Leu Lys Gln His Ala Ala Val Ser Glu Lys Leu Leu	
970 975 980	
tct ctt cta cca gag tat gtt gtt cca tat aca att cac ctt ttg gca	3149
Ser Leu Leu Pro Glu Tyr Val Val Pro Tyr Thr Ile His Leu Leu Ala	
985 990 995 1000	
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His Asp Pro Asp Tyr Val Lys Val Gln Asp Ile Glu Gln Leu Lys Asp	
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gtt aaa gaa tgt ctt tgg ttt gtt ctg gaa ata tta atg gct aaa aat	3245
Val Lys Glu Cys Leu Trp Phe Val Leu Glu Ile Leu Met Ala Lys Asn	
1020 1025 1030	
gaa aat aac agt cac gct ttt atc aga aag atg gta gaa aat att aaa	3293
Glu Asn Asn Ser His Ala Phe Ile Arg Lys Met Val Glu Asn Ile Lys	
1035 1040 1045	
caa aca aaa gat gcc caa gga cca gat gat gca aaa atg aat gaa aaa	3341
Gln Thr Lys Asp Ala Gln Gly Pro Asp Asp Ala Lys Met Asn Glu Lys	
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ctg tac act gtg tgt gat gtt gcc atg aat atc atc atg tca aag agt	3389
Leu Tyr Thr Val Cys Asp Val Ala Met Asn Ile Ile Met Ser Lys Ser	
1065 1070 1075 1080	
act aca tac agt ttg gaa tct cct aaa gac ccg gta cta cca gct cgt	3437
Thr Thr Tyr Ser Leu Glu Ser Pro Lys Asp Pro Val Leu Pro Ala Arg	
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ttc ttc act caa cct gac aag aat ttc agt aac acc aaa aat tat ctg	3485
Phe Phe Thr Gln Pro Asp Lys Asn Phe Ser Asn Thr Lys Asn Tyr Leu	
1100 1105 1110	
cct cct gaa atg aaa tca ttt ttc act cct gga aaa cct aaa aca acc	3533
Pro Pro Glu Met Lys Ser Phe Phe Thr Pro Gly Lys Pro Lys Thr Thr	
1115 1120 1125	
aat gtt cta gga gct gtt aac aag cca ctt tca tca gca ggc aag caa	3581
Asn Val Leu Gly Ala Val Asn Lys Pro Leu Ser Ser Ala Gly Lys Gln	
1130 1135 1140	
tct cag acc aaa tca tca cga atg gaa act gta agc aat gca agc agc	3629
Ser Gln Thr Lys Ser Ser Arg Met Glu Thr Val Ser Asn Ala Ser Ser	
1145 1150 1155 1160	

agc tca aat cca agc tct cct gga aga ata aag ggg agg ctt gat agt	3677
Ser Ser Asn Pro Ser Ser Pro Gly Arg Ile Lys Gly Arg Leu Asp Ser	
1165 1170 1175	
tct gaa atg gat cac agt gaa aat gaa gat tac aca atg tct tca cct	3725
Ser Glu Met Asp His Ser Glu Asn Glu Asp Tyr Thr Met Ser Ser Pro	
1180 1185 1190	
ttg ccg ggg aaa aaa agt gac aag aga gac gac tct gat ctt gta agg	3773
Leu Pro Gly Lys Lys Ser Asp Lys Arg Asp Asp Ser Asp Leu Val Arg	
1195 1200 1205	
tct gaa ttg gag aag cct aga ggc agg aaa aaa acg ccc gtc aca gaa	3821
Ser Glu Leu Glu Lys Pro Arg Gly Arg Lys Lys Thr Pro Val Thr Glu	
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cag gag gag aaa tta ggt atg gat gac ttg act aag ttg gta cag gaa	3869
Gln Glu Glu Lys Leu Gly Met Asp Asp Leu Thr Lys Leu Val Gln Glu	
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cag aaa cct aaa ggc agt cag cga agt cgg aaa aga ggc cat acg gct	3917
Gln Lys Pro Lys Gly Ser Gln Arg Ser Arg Lys Arg Gly His Thr Ala	
1245 1250 1255	
tca gaa tct gat gaa cag cag tgg cct gag gaa aag agg ctc aaa gaa	3965
Ser Glu Ser Asp Glu Gln Gln Trp Pro Glu Glu Lys Arg Leu Lys Glu	
1260 1265 1270	
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Asp Ile Leu Glu Asn Glu Asp Glu Gln Asn Ser Pro Pro Lys Lys Gly	
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aaa aga ggc cga cca cca aaa cct ctt ggt gga ggt aca cca aaa gaa	4061
Lys Arg Gly Arg Pro Pro Lys Pro Leu Gly Gly Gly Thr Pro Lys Glu	
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Glu Pro Thr Met Lys Thr Ser Lys Lys Gly Ser Lys Lys Lys Ser Gly	
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cct cca gca cca gag gag gag gaa gaa gaa gaa aga caa agt gga aat	4157
Pro Pro Ala Pro Glu Glu Glu Glu Glu Glu Arg Gln Ser Gly Asn	
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acg gaa cag aag tcc aaa agc aaa cag cac cga gtg tca agg aga gca	4205
Thr Glu Gln Lys Ser Lys Ser Lys Gln His Arg Val Ser Arg Arg Ala	
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Gln Gln Arg Ala Glu Ser Pro Glu Ser Ser Ala Ile Glu Ser Thr Gln	
1355 1360 1365	
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Ser Thr Pro Gln Lys Gly Arg Gly Arg Pro Ser Lys Thr Pro Ser Pro	
1370 1375 1380	

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 Ser Gln Pro Lys Lys Asn Val
 1385 1390

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